



Bioinformatics Core Support for the Frederick National Laboratory for Cancer Research and NCI

Eric Stahlberg

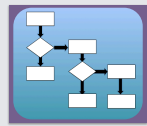
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Frederick
National
Laboratory
for Cancer Research

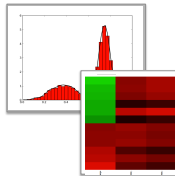


Exploring Experimental Options

- Exploration of new protocols
- Support for new technologies
- Custom training
- Bioinformatics consulting
- Establishing collaborations



Experimental Design



- Technology selection
- Feasibility analysis
- Statistical validation
- Study design
- Cost-saving recommendations
- Collaborations



Visit <http://ccrifx.cancer.gov> to get started

Sequence Data Request

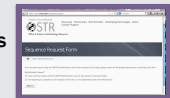


- Coordinated sequencing and analysis

Step 1: Submit analysis request at <http://ccrifx.cancer.gov>
Step 2: Submit sequencing request

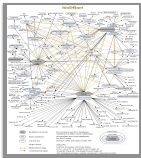
- Sequencing with self-analysis

Step 1. Submit sequencing request
Step 2. Coordinate with SF staff directly



Submit at <https://ostr.cancer.gov/sequence/request>

Hypothesis and Discovery



- Scientific insights
- Consulting on new techniques and emerging technologies
- Customized solutions
- In-house integration and annotation applications
- Guidance on future steps

ABCC bioinformatics efforts enable the NCI community by integrating community data, delivering key services, developing capabilities, building collaborations, and providing scientific consultation.

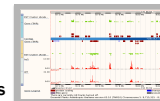


SAMPLE TO SCIENCE

*Collaborations, Core and Infrastructure
Enabling Cancer Research Advances*

Data Analysis

- Established NGS workflows
- ChIP-seq, RNA-seq, Exome-Seq
- Microarray analysis
- Data integration and pathway analysis
- Recognized scientific contributions in major journals
- CCRIFX analysis at no-cost to CCR investigators
- Collaborative analysis with over 200 scientists
- Over 100 projects completed



<http://ccrifx.cancer.gov>

Experimentation



The SF and LMT sequencing facilities provide sequencing support for Illumina, Ion Torrent, Pacific Biosciences, Roche/454



- More than 50 labs served
- Over 1300 sample lanes run

<http://ncifrederick.cancer.gov/atp/genetics-and-genomics/>

Results Integration and Delivery

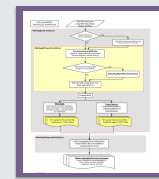
- Extended analysis with custom data integration
- Comparisons to existing datasets such as TCGA
- Custom, variation, impact and pedigree analysis
- Follow-up inquiries
- Publication support
- Custom visualizations
- Analysis data management



Data Quality Management

CCRFX and CCR SF staff coordinate to assess data quality, deliver original data and get the most from your experimental investment.

- Robust QC workflows
- Online quality reports



Visit <http://isp.ncifcrf.gov/abcc> for more information on these and other bioinformatics opportunities

